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SEQUENCE LISTING

<110> Bristol-Myers Squibb Company

<120> POLYNUCLEOTIDES ENCODING A NOVEL TESTIS-SPECIFIC TUBULIN
TYROSINE-LIGASE-LIKE PROTEIN, BGS42

<130> D0283A CIP

<150> U.S. 60/394,725
<151> 2002-07-09

<150> U.S.10/615,659
<151> 2003-07-09

<160> 103

<170> PatentIn version 3.2

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Tyr Phe Ser Gln Cys Gln Ala Leu Leu Asn Arg Ile Thr Ser Val Asn
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Glu Ile Leu Glu Leu Ala Ala Ala Asp His Pro Leu Ser Arg Asp Asn
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Ala Val Gln Lys Tyr Leu Lys Asn Asp Val Gly Arg Ser Pro Leu Leu
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Arg	Gly	Arg	Asp	Ile	Val	Cys	Met	Asp	Arg	Val	Glu	Glu	Ile	Leu	Glu		
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Ile Arg Gln Trp Phe Leu Val Thr Asp Trp Asn Pro Leu Thr Ile Trp
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Phe Tyr Lys Glu Ser Tyr Leu Arg Phe Ser Thr Gln Arg Phe Ser Leu
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Asp Lys Leu Asp Ser Ala Ile His Leu Cys Asn Asn Ala Val Gln Lys
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Lys Asn Leu Asp Asn Ser Val His Leu Cys Asn Asn Ser Ile Gln Lys
 100 105 110

His Leu Glu Asn Ser Cys His Arg His Pro Leu Leu Pro Pro Asp Asn
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Val Val Ile Asp Arg Met Leu Asp Arg Asn Cys Asp Thr Gly Ala Phe
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Glu Leu Ile Tyr Lys Gln Pro Val Thr Thr Ser Pro Ala Ser Thr Pro
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Ile Lys Thr Ser Pro Glu Leu Ala Glu Ser Cys Thr Trp Phe Pro Glu
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Ser Tyr Val Ile Tyr Pro Thr Asn Leu Lys Thr Pro Val Ala Pro Ala
100 105 110

Gln Asn Gly Ile His Pro Pro Ile His Ser Ser Arg Thr Asp Glu Arg
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Glu Phe Phe Leu Thr Ser Tyr Asn Lys Lys Lys Glu Asp Gly Glu Gly
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Asn Val Trp Ile Ala Lys Ser Ser Ala Gly Ala Lys Gly Glu Gly Ile
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Leu Ile Ser Ser Glu Ala Thr Glu Leu Leu Asp Phe Ile Asp Asn Gln
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Gly Gln Val His Val Ile Gln Lys Tyr Leu Glu Arg Pro Leu Leu Leu
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Glu Pro Gly His Arg Lys Phe Asp Ile Arg Ser Trp Val Leu Val Asp
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Ser Glu Pro Tyr His Thr Asp Asn Phe Gln Asp Lys Thr Cys His Leu
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Thr Asn His Cys Ile Gln Lys Glu Tyr Ser Lys Asn Tyr Gly Lys Tyr
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Glu Glu Gly Asn Glu Met Phe Phe Glu Glu Phe Asn Gln Tyr Leu Thr
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Ala Cys Ala Gln Lys Leu Tyr Ala Glu Leu Cys Gln Gly Ile Val Asp
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Arg Asn Ile Trp Ile Val Lys Pro Gly Ala Lys Ser Arg Gly Arg Gly
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Asn Pro Val Val Met Lys Asp Gly Lys Trp Val Val Gln Lys Tyr Ile
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Glu Arg Pro Leu Leu Ile Phe Gly Thr Lys Phe Asp Leu Arg Gln Trp
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Phe Leu Val Thr Asp Trp Asn Pro Leu Thr Val Trp Phe Tyr Arg Asp
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Asn Ser Val His Leu Cys Asn Asn Ser Ile Gln Lys His Leu Glu Asn
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Tyr Gly Ala Asp Phe Val Phe Gly Glu Asp Phe Gln Pro Trp Leu Ile
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Arg Leu Cys Ala Gly Val Gln Ala Asp Thr Leu Arg Val Val Ile Asp
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Leu Leu Pro Met Tyr Ser Asp Thr Arg Ala Arg Ser Ser Asp Asp Ser

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Lys	Trp	Val	Val	Gln	Lys	Tyr	Ile	Glu	Thr	Pro	Leu	Leu	Ile	Cys	Asp
			180					185						190	
Thr	Lys	Phe	Asp	Ile	Arg	Gln	Trp	Phe	Leu	Val	Thr	Asp	Trp	Asn	Pro

195	200	205
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Arg Phe Ser Leu Asp Lys Leu Asp Ser Ala Ile His Leu Cys Asn Asn 225 230 235 240		
Ala Val Gln Lys Tyr Leu Lys Asn Asp Val Gly Arg Ser Pro Leu Leu 245 250 255		
Pro Ala His Asn Met Trp Thr Ser Thr Arg Phe Gln Glu Tyr Leu Gln 260 265 270		
Arg Gln Gly Arg Gly Ala Val Trp Gly Ser Val Ile Tyr Pro Ser Met 275 280 285		
Lys Lys Ala Ile Ala His Ala Met Lys Val Ala Gln Asp His Val Glu 290 295 300		
Pro Arg Lys Asn Ser Phe Glu Leu Tyr Gly Ala Asp Phe Val Leu Gly 305 310 315 320		
Arg Asp Phe Arg Pro Trp Leu Ile Glu Ile Asn Ser Ser Pro Thr Met 325 330 335		
His Pro Ser Thr Pro Val Thr Ala Gln Leu Cys Ala Gln Val Gln Glu 340 345 350		
Asp Thr Ile Lys Val Ala Val Asp Arg Ser Cys Asp Ile Gly Asn Phe 355 360 365		
Glu Leu Leu Trp Arg Gln Pro Val Val Glu Pro Pro Pro Phe Ser Gly 370 375 380		
Ser Asp Leu Cys Val Ala Gly Val Ser Val Arg Arg Ala Arg Arg Gln 385 390 395 400		
Val Leu Pro Val Cys Asn Leu Lys Ala Ser Ala Ser Leu Leu Asp Ala 405 410 415		
Gln Pro Leu Lys Ala Arg Gly Pro Ser Ala Met Pro Asp Pro Ala Gln 420 425 430		

Gly Pro Pro Ser Pro Ala Leu Gln Arg Asp Leu Gly Leu Lys Glu Glu
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Lys Gly Leu Pro Leu Ala Leu Leu Ala Pro Leu Arg Gly Ala Ala Glu
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Ser Gly Gly Ala Ala Gln Pro Thr Arg Thr Lys Ala Ala Gly Lys Val
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Glu Leu Pro Ala Cys Pro Cys Arg His Val Asp Ser Gln Ala Pro Asn
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Thr Gly Val Pro Val Ala Gln Pro Ala Lys Ser Trp Asp Pro Asn Gln
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 35 40 45

Leu Asn Arg Ile Thr Ser Val Asn Pro Gln Thr Asp Ile Asp Gly Leu
 50 55 60

Arg Asn Ile Trp Ile Ile Lys Pro Ala Ala Lys Ser Arg Gly Arg Asp
 65 70 75 80

Ile Val Cys Met Asp Arg Val Glu Glu Ile Leu Glu Leu Ala Ala Ala
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Asp His Pro Leu Ser Arg Asp Asn Lys Trp Val Val Gln Lys Tyr Ile
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Glu Thr Pro Leu Leu Ile Cys Asp Thr Lys Phe Asp Ile Arg Gln Trp
 115 120 125

Phe Leu Val Thr Asp Trp Asn Pro Leu Thr Ile Trp Phe Tyr Lys Glu
 130 135 140

Ser Tyr Leu Arg Phe Ser Thr Gln Arg Phe Ser Leu Asp Lys Leu Asp
 145 150 155 160

Ser Ala Ile His Leu Cys Asn Asn Ala Val Gln Lys Tyr Leu Lys Asn
 165 170 175

Asp Val Gly Arg Ser Pro Leu Leu Pro Ala His Asn Met Trp Thr Ser
 180 185 190

Thr Arg Phe Gln Glu Tyr Leu Gln Arg Gln Gly Arg Gly Ala Val Trp
 195 200 205

Gly Ser Val Ile Tyr Pro Ser Met Lys Lys Ala Ile Ala His Ala Met
 210 215 220

Lys Val Ala Gln Asp His Val Glu Pro Arg Lys Asn Ser Phe Glu Leu
 225 230 235 240

Tyr Gly Ala Asp Phe Val Leu Gly Arg Asp Phe Arg Pro Trp Leu Ile
 245 250 255

Glu Ile Asn Ser Ser Pro Thr Met His Pro Ser Thr Pro Val Thr Ala
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Arg Ser Cys Asp Ile
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Asn

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 35 40 45
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 Asp Tyr Phe Glu Lys Arg Glu Arg Glu Gly Gln Asp Asn Thr Trp Ile
 65 70 75 80
 Val Lys Pro Ser Ala Ser Ala Arg Gly Arg Gly Ile Arg Ile Thr Asn
 85 90 95
 Lys Leu Ser Gln Ile Pro Lys Trp Val Asp Asp Phe Lys Asp Asn Pro
 100 105 110
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 Asp Leu Arg Leu Tyr Val Leu Val Thr Ser Val Asp Pro Leu Arg Ile
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 Tyr Val Tyr Arg Glu Gly Leu Leu Arg Phe Ala Ser Val Lys Tyr Ser
 165 170 175
 Pro Leu Ser Asp Leu Asp Asp Val Glu Met His Leu Thr Asn Tyr Ser
 180 185 190
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 Glu Pro Val Gly His Lys Trp Ser Leu Gln Asn Leu Trp Leu Tyr Leu
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 Glu Glu Met Gly Ile Asp Lys Glu Glu Ile Trp Leu Gln Ile Glu Ser

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Ile Met Leu Lys Trp Asp Asp Asp Glu Asn Leu Lys Pro Trp Leu Leu						
	275		280		285	
Glu Val Asn Ala Ser Pro Ser Leu His Ser Thr Thr Lys Leu Asp Ala						
	290		295		300	
Glu Leu Lys Ala Pro Leu Ile Arg Asp Val Leu Asn Leu Ala Leu Ser						
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Tyr Glu Leu His Tyr Met Glu Met Gly Asp Leu Tyr Leu Ile Asn Gly
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Glu Ala Arg Ala His Thr Arg Thr Leu Asn Val Lys Gln Asn Tyr Glu
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Glu Trp Phe Ser Phe Val Gly Glu Gln Asp Leu Pro Leu Ala Asp Leu
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Asp Val Ile Leu Met Arg Lys Asp Pro Pro Phe Asp Thr Glu Phe Ile
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Tyr Ala Thr Tyr Ile Leu Glu Arg Ala Glu Glu Lys Gly Thr Leu Ile
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Val Asn Lys Pro Gln Ser Leu Arg Asp Cys Asn Glu Lys Leu Phe Thr
 115 120 125

Ala Trp Phe Ser Asp Leu Thr Pro Glu Thr Leu Val Thr Arg Asn Lys
 130 135 140

Ala Gln Leu Lys Ala Phe Trp Glu Lys His Ser Asp Ile Ile Leu Lys
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Pro Leu Asp Gly Met Gly Gly Ala Ser Ile Phe Arg Val Lys Glu Gly
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Asp Pro Asn Leu Gly Val Ile Ala Glu Thr Leu Thr Glu His Gly Thr
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Arg Tyr Cys Met Ala Gln Asn Tyr Leu Pro Ala Ile Lys Asp Gly Asp
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Lys Arg Val Leu Val Val Asp Gly Glu Pro Val Pro Tyr Cys Leu Ala
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Arg Ile Pro Gln Gly Gly Glu Thr Arg Gly Asn Leu Ala Ala Gly Gly
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Asp Ile Ile Gly Asp Arg Leu Thr Glu Ile Asn Val Thr Ser Pro Thr
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Cys Ile Arg Glu Ile Glu Ala Glu Phe Pro Val Ser Ile Thr Gly Met
290 295 300

Leu Met Asp Ala Ile Glu Ala Arg Leu Gln
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